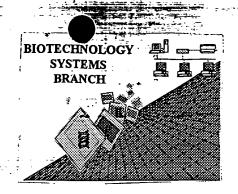
RAW-SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

JAN 05 7001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS. PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION OUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2Kcompliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

Raw Sequence Listing Error Summary

ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/497,997

Wrapped Nucleis The numberlext at the end of each line "wapped" down to the next Ene. This may occur by four file was retireved in a word processor after creating it. Please adjust your right margin to 3, as this will prevent "wrapping". 2 Wrapped Aminos The amino acid numberriext at the end of each line "wrapped" down to the next line. This may occur your file was retrieved in a word processor after creating it. Please adjust your right margin to 3, as this will prevent "wrapping". 3 Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces. 4 Misaligned Amino Acid The numbering under each 5th amino acid is mijalagned. This may be caused by the use of labs between the numbering. It is recommended to delete any tabs and use spacing between the numbers. 5 Non-ASCII This file was not saved in ASCII (DOS) lord, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII total so that it can be processed. 6 Variable Length Sequence(s) contain n's or Xaa's which represented more than one residue. As per the rules, each nor Xaa can only represent a single recidue. Please present the maximum number of each residue having variable length and indicate in the (of feature section that some may be missing. 7 Patentin ver. 2.0 "bug" Sequence(s) contain version 2.0 has caused the *220" *222" section to be missing from amino acid sequences (out the section of the previously coded nucleic acid sequence. Please manually copy the retevant *220" *222" section from the previously coded nucleic acid sequence. Please manually copy the retevant *220" *222" sections for Artificial or furknown sequences. 8 Skipped Sequences (a) International, please use the following format for each skipped sequence (2) Sequence(s) missing if intentional, please use the following format for each skipped sequence (3) Sequence is intentionally skipped Please astos adjust the "(a) NUMBER OF SEQUENCES" response to includ	ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE			
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13 Patentln ver. 2.0 "bug" Please do not use "Copy to Disk" function of Patentln version 2.0. This causes a corrupted) (
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AKS-Biotechnology Systems Branch- 5/15/99

Instead, please use "File Manager" or any other means to copy file to floppy disk.

1648 Diskette Net RAW SEQUENCE LISTING DATE: 12/26/2000 PATENT APPLICATION: US/09/497,997 TIME: 14:24:34 Corrected Diskette Needed Input. Set : A:\06600166.txt Output Set: N:\CRF3\12262000\1497997.raw 3 <110> APPLICANT: TERNYNCK, THERESA AVRAMEAS, ALEXANDRE BUTTIN, GERARD AVRAMEAS, STRAITTS SARON, MARIE-FRANCOISE BLONDEL, BRUNO 8 COUDERC, THERESA 10 MICHELSON, SUSAN 1.1 ZIPETO, DONATO 13 <120> TITLE OF INVENTION: VECTORS DERIVED FROM ANTIBODIES FOR TRANSFERRING SUBSTANCES INTO CELLS 15 <130> FILE REFERENCE: 0660-0166-0XCONT 17 <140> CURRENT APPLICATION NUMBER: 09/497,997 18 <141> CURRENT FILING DATE: 2000-02-04 20 <150> PRIOR APPLICATION NUMBER: PCT/FR98/01740 21 <151> PRTOR FILING DATE: 1998-08-04 Ker 1.823 d) nen Seguera Rules, the only valid (2137 responses are: Unknown, 23 <160> NUMBER OF SEO ID NOS: 35 25 <170> SOFTWARE: PatentIn version 3.0 27 <210> SEQ ID NO: 1 28 <211> LENGTH: 17 29 <212> TYPE: PRT Artificial/Unknown 30 <213> ORGANISM Artificial Sequence, or scientific name 32 <220> FEATURE: 33 <221> NAME/KEY: misc_feature 34 <222> LOCATION: ()..() (benus/species) 35 <223> OTHER INFORMATION: Description of Artificial Sequence: peptide 38 <400> SEQUENCE: 1 40 Thr Arg Gln Lys Tyr Asn Lys Arg Ala Met Asp Tyr Trp Gly Gln Gly 41 1 43 Thr 46 <210> SEQ ID NO: 2 47 <211> LENGTH: 17 48 <212> TYPE: PRT 49 <213> ORGANISM Artificial/Unknown 51 <220> FEATURE: 52 <221> NAME/KEY: misc_feature 53 <222> LOCATION: ()..() 54 <223> OTHER INFORMATION: Description of Artificial Sequence: peptide 57 <400> SEQUENCE: 2 give source of senetic material-see circled portion of them 12 on Ever Summar Sheet 59 Thr Arg Gln Lys Tyr Gly Lys Arg Gly Met Asp Tyr Trp Gly Gln Gly 60 1 62 Thr 65 <210> SEQ ID NO: 3 66 <21.1> LENGTH: 18 67 <212> TYPE: PRT. 68 <213> ORGANISM: Artificial/Unknown 70 <220> FEATURE: 71 <221> NAME/KEY: misc_feature

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PATENT APPLICATION: US/09/497,997

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